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January 11, 2005, 18:45:51; Search time 466 Seconds (without alignments) 35.665 Million cell updates/sec

US-09-733-179A-2 84 Title: Perfect score:

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6730630 seqs, 1107998698 residues Searched:

6730630 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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